

Wang

CRF Errors Corrected by the STIC System Branch

1635

11/5/98

CRF Processing Date: _____
Edited by: _____
Verified by: _____ (STIC staff)

Serial Number: 08/978,636

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: **ENTERED**
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file;
☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/978,636DATE: 11/05/98
TIME: 14:41:32

INPUT SET: S29661.raw

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

SEQUENCE LISTING

1
2
3 (1) General Information
4
5 (i) APPLICANT: ELAZAR RABBANI
6 JANNIS G. STAVRIANOPOULOS
7 JAMES J. DONEGAN
8 DAKAI LIU
9 NORMAN E. KELKER
10 DEAN L. ENGELHARDT
11
12 (ii) TITLE OF INVENTION: NOVEL PROPERTY EFFECTING
13 AND/OR PROPERTY EXHIBITING COMPOSITIONS FOR
14 THERAPEUTIC AND DIAGNOSTIC USE
15
16 (iii) NUMBER OF SEQUENCES: 42
17
18 (iv) CORRESPONDENCE ADDRESS:
19 (A) ADDRESSEE: ENZO THERAPEUTICS, INC.
20 (B) STREET: C/O ENZO BIOCHEM, INC.
21 527 MADISON AVENUE,
22 9TH FLOOR
23 (C) CITY: NEW YORK
24 (D) STATE: NY
25 (E) COUNTRY: USA
26 (F) ZIP: 10022
27
28 (v) COMPUTER READABLE FORM:
29 (A) MEDIUM TYPE: 3.5" Micro Floppy Disk. 1.44 KB
30 STORAGE
31 (B) COMPUTER: IBM PC/XT/AT, IBM PS/2 OR COMPATIBLES
32 (C) OPERATING SYSTEM: PC-DOS
33 (D) SOFTWARE: MICROSOFT WORD ____ - ASCII TEXT (DOS)
34
35 (vi) CURRENT APPLICATION DATA:
36 (A) APPLICATION NUMBER: US 08/978,636
37 (B) FILING DATE: 25-NOVEMBER-1997
38 (C) CLASSIFICATION: Not Yet Known
39
40 (viii) ATTORNEY/AGENT INFORMATION:
41 (A) NAME: FEDUS, RONALD C.
42 (B) REGISTRATION NUMBER: 32,567
43 (C) REFERENCE/DOCKET NUMBER: ENZ-53(D3)
44
45 (ix) TELECOMMUNICATION INFORMATION:

RAW SEQUENCE LISTING
PATENT APPLICATION *US/08/978,636*DATE: 11/05/98
TIME: 14:41:33*INPUT SET: S29661.raw*

46 (A) TELEPHONE: (212) 583-0100
47 (B) TELEFAX: (212) 583-0150
48

ERRORED SEQUENCES FOLLOW:

814 (2) INFORMATION FOR SEQ ID NO:42:
815
816 (i) SEQUENCE CHARACTERISTICS:
--> 817 (A) **LENGTH:67 base pairs**
818 (B) TYPE:nucleic acid
819 (C) STRANDEDNESS:single
820 (D) TOPOLOGY:linear
821
822 (ii) MOLECULE TYPE: other nucleic acid
823 (A) DESCRIPTION: /desc = "oligonucleotide"
824
825 (iii) HYPOTHETICAL: YES
826
827 (ix) SEQUENCE DESCRIPTION:SEQ ID NO:42:
828
829 CCGGATAATA CGACTCACTA TAGGGCGAGC TCGGTACCCG GGTCTAGAGT CGACCTGCAG 60
830
831 GCATGCT 67
832
833
--> 834 **SEQ ID: PAGE 16**
835
836

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/978,636

DATE: 11/05/98
TIME: 14:41:34

INPUT SET: S29661.raw

Line	Error	Original Text
38	Wrong Classification	(C)CLASSIFICATION:Not Yet Known
817	Entered (67) and Calc. Seq. Length (71) differ	(A)LENGTH:67 base pairs
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	# of Sequences for line conflicts w/ running total	SEQ ID: PAGE 16